



1600

RAW SEQUENCE LISTING

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:21:50

Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

4 <110> APPLICANT: Massague et al.
 6 <120> TITLE OF INVENTION: ISOLATED p27 PROTEIN AND METHODS FOR ITS PRODUCTION AND USE
 8 <130> FILE REFERENCE: GPC1-P03-079
 10 <140> CURRENT APPLICATION NUMBER: 09/865018B
 11 <141> CURRENT FILING DATE: 2001-05-24
 13 <150> PRIOR APPLICATION NUMBER: 08/854039
 14 <151> PRIOR FILING DATE: 1997-05-09
 16 <160> NUMBER OF SEQ ID NOS: 27
 18 <170> SOFTWARE: Patentix version 3.1
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 597
 22 <212> TYPE: DNA
 23 <213> ORGANISM: Homo sapiens
 24 <210> FEATURE:
 25 <211> NAME/KEY: CDS
 26 <212> LOCATION: (1)..(597)
 27 <213> OTHER INFORMATION:

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52	Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met	
53	5 10 15	
54	acc gcc agg cag gcc gag cac ccc aag ccc tgg gcc tgc agg aac ctg	96
55	Asp Ala Arg Gln Ala Glu His Pro Lys Pro Ser Ala Cys Arg Asn Leu	
56	20 25 30	
57	tcc ggt cgg ggc gac cac gaa gag tta acc cgg gac ttg gag aag cac	144
58	Phe Gly Pro Val Asp His Glu Gln Leu Thr Arg Asp Leu Glu Lys His	
59	35 40 45	
60	tgc aga gac atg caa gag gcc agc cag cgc aag tgg aat ttc gat ttt	192
61	Cys Arg Asp Met Gln Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe	
62	50 55 60	
63	caa aat cac aaa ccc cta gag gcc aag tac gag tgg caa gag gtg gag	240
64	Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu	
65	65 70 75 80	
66	aag gcc agc ttg ccc gag ttc tac tac aga ccc ccg cgg ccc ccc aaa	288
67	Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys	
68	85 90 95	
69	ggc gcc tgc aag ctg ccg gcc caa gag agc caa gat gtc agc cgg agc	336
70	Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Ser	
71	100 105 110	
72	cgc ccg gcc gcc cct tta att gga gct ccg gct aac tct gag gac acc	384
73	Arg Pro Ala Ala Pro Leu Ile Gly Ala Pro Ala Asn Ser Glu Asp Thr	
74	115 120 125	
75	cac ttg gtg gac caa aag act gat ccg tgg gac acc cag aac gga tta	432

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Output Set: N:\CRF4\04232003\I865018B.raw

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64 His Leu Val Asp Pro Lys Thr Asp Pro Ser Asp Ser Gln Thr Gly Leu
65      130      135      140
67 ggc gag caa ttc gca gga ata agg aag oga oot gca acc gac gat tat      480
68 Ala Glu Gln Cys Ala Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
69 145      150      155      160
71 tct act caa acc ata aga gcc aac aga aca gaa gaa aat gtt tca gat      528
72 Ser Thr Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
73      165      170      175
75 ggt tcc cca aat gac ggt tct gtg gag cag aag ccc aag aag cct ggt      576
76 Gly Ser Pro Asn Ala Gly Ser Val Glu Gln Thr Pro Lys Lys Pro Gly
77      180      185      190
79 ctc aga aga cgt cca aag taa      597
80 Leu Arg Arg Arg Gln Thr
81      195
83 <210> SEQ ID NO: 2
84 <211> LENGTH: 198
85 <212> TYPE: PRT
86 <213> ORGANISM: Homo sapiens
87 <400> SEQUENCE: 2
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90      5      10      15
94 Asp Ala Arg Gln Ala Glu His Pro Lys Pro Ser Ala Cys Arg Asn Leu
95      20      25      30
98 Phe Gly Pro Val Asp His Glu Glu Leu Thr Arg Asp Leu Glu Lys His
99      35      40      45
102 Cys Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
103      50      55      60
106 Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu
107 65      70      75      80
110 Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
111      85      90      95
114 Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Ser
115      100      105      110
118 Arg Pro Ala Ala Pro Leu Ile Gly Ala Pro Ala Asn Ser Glu Asp Thr
119      115      120      125
122 His Leu Val Asp Pro Lys Thr Asp Pro Ser Asp Ser Gln Thr Gly Leu
123      130      135      140
126 Ala Glu Gln Cys Ala Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
127 145      150      155      160
130 Ser Thr Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
131      165      170      175
134 Gly Ser Pro Asn Ala Gly Ser Val Glu Gln Thr Pro Lys Lys Pro Gly
135      180      185      190
138 Leu Arg Arg Arg Gln Thr
139      195
141 <210> SEQ ID NO: 3
142 <211> LENGTH: 594
143 <212> TYPE: DNA
144 <213> ORGANISM: Mus musculus

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Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

146 <1070> FEATURE:

147 <1071> NAME/KEY: CDS

148 <1072> LOCATION: (1)..(594)

149 <1073> OTHER INFORMATION:

W--> 151 <400> 3

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153 1          5          10          15
154 gac gcc aga caa gag gat cac ccc aag cct tcc gcc tgc aga aat ctg      96
155 Asp Ala Arg Gln Ala Asp His Pro Lys Pro Ser Ala Cys Arg Asn Leu
156 20          25          30
157 ttc ggc cag gtc aat cat gaa gaa cta acc cgg gac ttg gag aag cac      144
158 Phe Gly Pro Val Asn His Glu Glu Leu Thr Arg Asp Leu Glu Lys His
159 35          40          45
160 tcc cgg gat atg gaa gaa gag agt cag cgc aag tgg aat ttc gac ttt      192
161 Cys Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
162 50          55          60
163 cag aat cat aag ccc ctg gag ggc aga tac gaa tgg cag gag gtg gag      240
164 Gln Asn His Lys Pro Leu Glu Gly Arg Tyr Glu Trp Gln Glu Val Glu
165 65          70          75          80
166 aac gcc agc ttc ccc gag ttc tac tac agg ccc cag cgc ccc ccc aag      288
167 Arg Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
168 85          90          95
169 gac ccc tgc aag ggc ctg ggc cag gag agc cag gat gtc agc ggg agc      336
170 Ser Ala Cys Lys Val Leu Ala Gln Glu Ser Gln Asp Val Ser Gly Ser
171 100          105          110
172 cgc cag gag gtg cct tta att ggg tot cag gca aac tot gag gac cgg      384
173 Arg Gln Ala Val Pro Leu Ile Gly Ser Gln Ala Asn Ser Glu Asp Arg
174 115          120          125
175 cat ttg gtg gac caa atg cct gac tgg tca gac aat cag got ggg tta      432
176 His Leu Val Asp Gln Met Pro Asp Ser Ser Asp Asn Gln Ala Gly Leu
177 130          135          140
178 gag cag cag tot cca ggg atg agg aag cga cct got gca gaa gat tot      480
179 Ala Glu Gln Cys Pro Gly Met Arg Lys Arg Pro Ala Ala Glu Asp Ser
180 145          150          155          160
181 tct tgg caa aac aaa agg gcc aac aga aca gaa gaa aat gtt tca gac      528
182 Ser Ser Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
183 165          170          175
184 cat tcc cag aac got ggc act gtg gag cag acg ccc aag aag ccc ggc      576
185 Gly Ser Pro Asn Ala Gly Thr Val Glu Gln Thr Pro Lys Lys Pro Gly
186 180          185          190

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200 att cga cgc cag acg taa

594

201 Leu Arg Arg Gln Thr

202 195

203 <100> SEQ ID NO: 4

204 <110> LENGTH: 195

205 <120> TYPE: CDS

206 <130> ORGANISM: Mus musculus

207 <400> SEQUENCE: 4

RAW SEQUENCE LISTING

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TIME: 07:21:50

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Output Set: N:\CRF4\04232003\I865018B.raw

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211 Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met
212 1 10 15
215 Asp Ala Arg Gln Ala Asp His Pro Lys Pro Ser Ala Cys Arg Asn Leu
216 20 25 30
219 Phe Gly Pro Val Asn His Glu Gln Leu Thr Arg Asp Leu Glu Lys His
220 35 40 45
223 Cys Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
224 50 55 60
227 Gln Asn His Lys Pro Leu Glu Gly Arg Tyr Glu Trp Gln Glu Val Glu
228 65 70 75 80
231 Arg Gly Ser Leu Pro Gln Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
232 85 90 95
235 Ser Ala Cys Lys Val Leu Ala Gln Glu Ser Gln Asp Val Ser Gly Ser
236 100 105 110
239 Arg Gln Ala Val Pro Leu Ile Gly Ser Gln Ala Asn Ser Glu Asp Arg
240 115 120 125
243 His Leu Val Asp Gln Met Pro Asp Ser Ser Asp Asn Gln Ala Gly Leu
244 130 135 140
247 Ala Gln Gln Cys Pro Gly Met Arg Lys Arg Pro Ala Ala Glu Asp Ser
248 145 150 155 160
251 Ser Ser Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
252 165 170 175
255 Gly Ser Pro Asn Ala Gly Thr Val Glu Gln Thr Pro Lys Lys Pro Gly
256 180 185 190
259 Leu Arg Arg Gln Thr
260 195

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261 <210> SEQ ID NO: 5

262 <211> LENGTH: 534

263 <212> TYPE: DNA

264 <213> ORGANISM: Mustela vison

265 <214> FEATURE:

266 <215> NAME/KEY: CDS

267 <216> LOCATION: (1)..(534)

268 <217> OTHER INFORMATION:

W--> 272 <400> 5

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273 att tca aac atg cgg gtg tct aac gag agc cgg aac atg gag cgg atg 48
274 Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met
275 1 10 15
278 gag gcc aga cag cgg gag tac ccc aag ccc tcc gcc tgc aga aac ctc 96
279 Asp Ala Arg Gln Ala Glu Tyr Pro Lys Pro Ser Ala Cys Arg Asn Leu
280 20 25 30
283 ttc cgc cgg ctc aac cac gaa gag ctg acc cgg gac ttg gag aag cac 144
284 Phe Gly Pro Val Asn His Glu Gln Leu Thr Arg Asp Leu Glu Lys His
285 35 40 45
288 ccc aga gac atg gaa gag gca agc cag cgc aag tgg aat ttc gat ttc 192
289 Arg Arg Asp Met Glu Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
290 50 55 60
293 cca aat cac aag ccc ctg gag ggc aaa tac gag tgg cag gag gtg gag 240
294 Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu

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RAW SEQUENCE LISTING

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:21:50

Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

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291 65          70          75          80
293 aug ggc agc ttg cgg gag ttc tac tac aga ccc cgg cgg cca ccc aac 288
294 Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
295          85          90          95
296 ggc ggc tgc atg ttg cgg ggg cag gag agc cag gac gtc agc ggc aac 336
297 Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Thr
298          100          105          110
301 cgg cag gcc gtc cct tta atg ggg tct cag gca aac tca gag ga aca 384
302 Arg Gln Ala Val Pro Leu Met Gly Ser Gln Ala Asn Ser Glu Asp Thr
303          115          120          125
305 cag ttg gta gac aac aag act gac aag ggg gac aac cag gct ggc tta 432
306 His Leu Val Asp Gln Lys Thr Asp Thr Ala Asp Asn Gln Ala Gly Leu
307          130          135          140
309 ggc aag cag ttc act ggg atc app aag cga cgg gcc aca gac gat ttc 480
310 Ala Glu Gln Cys Thr Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
311          145          150          155          160
313 cct cct aac aac aac aga gcc aac aga aca gac gac aat gtc tca gac 528
314 Ser Pro Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
315          165          170          175
317 ggt tcc
318 Gly Ser
321 3210 - SEQ ID NO: 6
322 3211 - LENGTH: 177
323 3212 - TYPE: PRT
324 3213 - ORGANISM: Mustela vison
325 3214 - SEQUENCE: 6
326 Met Ser Asn Val Arg Val Ser Asn Gly Ser Pro Ser Leu Glu Arg Met
327          5          10          15
328 Asp Ala Arg Gln Ala Glu Tyr Pro Lys Pro Ser Ala Cys Arg Asn Leu
329          20          25          30
330 Phe Gly Pro Val Asn His Glu Glu Leu Thr Arg Asp Leu Glu Lys His
331          35          40          45
332 Arg Arg Asp Met Gln Glu Ala Ser Gln Arg Lys Trp Asn Phe Asp Phe
333          50          55          60
334 Gln Asn His Lys Pro Leu Glu Gly Lys Tyr Glu Trp Gln Glu Val Glu
335          65          70          75          80
336 Lys Gly Ser Leu Pro Glu Phe Tyr Tyr Arg Pro Pro Arg Pro Pro Lys
337          85          90          95
338 Gly Ala Cys Lys Val Pro Ala Gln Glu Ser Gln Asp Val Ser Gly Thr
339          100          105          110
340 Arg Gln Ala Val Pro Leu Met Gly Ser Gln Ala Asn Ser Glu Asp Thr
341          115          120          125
342 His Leu Val Asp Gln Lys Thr Asp Thr Ala Asp Asn Gln Ala Gly Leu
343          130          135          140
344 Ala Glu Gln Cys Thr Gly Ile Arg Lys Arg Pro Ala Thr Asp Asp Ser
345          145          150          155          160
346 Ser Pro Gln Asn Lys Arg Ala Asn Arg Thr Glu Glu Asn Val Ser Asp
347          165          170          175
352 Gly Ser

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/865,018B

DATE: 04/23/2003
TIME: 07:21:51

Input Set : N:\efs\09865018\09865018A\PTOMS.txt
Output Set: N:\CRF4\04232003\I865018B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; N Pos. 3,9
Seq#:13; N Pos. 1,13,16,22
Seq#:14; N Pos. 6,9,12,15
Seq#:15; N Pos. 6,9,12
Seq#:16; N Pos. 1,7,10,13,16
Seq#:17; N Pos. 1,10,13,16
Seq#:18; N Pos. 3
Seq#:19; N Pos. 15
Seq#:21; N Pos. 1

VERIFICATION SUMMARY

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:21:51

Input Set : N:\efs\09865018\09865018A\PTOMS.txt

Output Set: N:\CRF4\04232003\I865018B.raw

L:30 M:258 W: Mandatory Feature missing, <203> Blank for SEQ#:1,Line#:28
L:151 M:258 W: Mandatory Feature missing, <220> Blank for SEQ#:3,Line#:149
L:272 M:258 W: Mandatory Feature missing, <220> Blank for SEQ#:5,Line#:270
L:442 M:341 W: (46) "r" or "Xaa" used, for SEQ ID#:12 after pcs.:0
L:459 M:341 W: (46) "r" or "Xaa" used, for SEQ ID#:13 after pcs.:0
L:466 M:283 W: Missing Blank Line separator, <220> field identifier
L:474 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pcs.:0
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pcs.:0
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pcs.:0
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pcs.:0
L:536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pcs.:0
L:554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pcs.:0
L:581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pcs.:0



Does Not Comply
Corrected Diskette Needed

1600

RAW SEQUENCE LISTING

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:17:28

Input Set : N:\efs\09865018\09865018A\GPCI-P08-079SubstituteSequence.txt

Output Set: N:\CRF4\04232003\I865018B.raw

4 <110> APPLICANT: Massague et al.
 6 <120> TITLE OF INVENTION: ISOLATED p27 PROTEIN AND METHODS FOR ITS PRODUCTION AND USE
 8 <130> FILE REFERENCE: GPCI-P08-079
 10 <140> CURRENT APPLICATION NUMBER: 09/865018B
 11 <141> CURRENT FILING DATE: 2001-05-24
 13 <150> PRIOR APPLICATION NUMBER: 08/854039
 14 <151> PRIOR FILING DATE: 1997-05-09
 16 <160> NUMBER OF SEQ ID NOS: 27
 18 <170> SOFTWARE: PatentIn version 3.1

ERRORED SEQUENCES

643 <210> SEQ ID NO: 25
 644 <211> LENGTH: 6
 645 <212> TYPE: PRT
 646 <213> ORGANISM: Homo sapiens
 648 <240> SEQUENCE: 25
 650 Leu Phe Gly Pro Val Asp

E--> 651 1 5

Realign amino acid numbering

VERIFICATION SUMMARY

DATE: 04/23/2003

PATENT APPLICATION: US/09/865,018B

TIME: 07:17:29

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Output Set: N:\CRF4\04232003\I865018B.raw

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L:151 M:158 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:149
L:272 M:158 W: Mandatory Feature missing, <223> Blank for SEQ#:5,Line#:270
L:441 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:465 M:183 W: Missing Blank Line separator, <220> field identifier
L:474 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:538 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:651 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:25

CRFI

CRF Errors Corrected by the STIC Systems Branch

CRF Processing Date

Edited by: M. SPENCER

Verified by: _____

RECEIVED

MAY 12 2003 (STIC staff)

TECH CENTER 1600:2900

ENTERED

Serial Number: 09/865,0183

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: Realigned amino acid numbering in
SEQ ID #25

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95